

1. A method of identifying a user of a mobile device, comprising:  
receiving a request from the mobile device to identify the user;  
retrieving a user identifier from a database;  
determining whether the user identifier is associated with the mobile device;  
if the user identifier is associated with the mobile device, identifying the user as the user associated with the user identifier;  
if the user identifier is not associated with the mobile device, identifying the user as an unknown user.

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interaction site and using a computer algorithm to predict a three-dimensional representation of said potential binding agent.

10. The method of claim 1, wherein said plurality of atomic coordinates are as set forth in Table 1.
- 5 11. The method of claim 1, wherein said potential binding agent is designed de novo.
12. A WW domain binding agent identified by the method of claim 1.
13. A method of identifying a WW domain binding agent, said method comprising:
  - 10 (a) defining an interaction site of a WW domain based on a plurality of atomic coordinates of said WW domain;
  - (b) modeling a potential binding agent that fits spatially into said interaction site;
  - (c) 15 determining the ability of said potential binding agent to compete with a WW domain substrate for said interaction site by contacting said potential binding agent with said WW domain in the presence of said WW domain substrate.

14. A method of identifying a WW domain binding agent, said method comprising:
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- (a) modeling a potential binding agent that fits spatially into an interaction site of a WW domain defined by a plurality of atomic coordinates of said WW domain;
  - (b) contacting said potential binding agent with said WW domain in the presence of a WW domain substrate; and
  - (c) determining the ability of said potential binding agent to compete with said WW domain substrate for binding to said WW domain.
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15. A method of identifying a WW domain binding agent, said method comprising:
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- (a) modeling a potential binding agent that fits spatially into an interaction site of a WW domain defined by a plurality of atomic coordinates of said WW domain; and
  - (b) determining the ability of said potential binding agent to compete with a WW domain substrate for said interaction site by contacting said potential binding agent with said WW domain in the presence of said WW domain substrate.
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16. A method of identifying a WW domain binding agent, said method comprising:
- determining the ability of a potential binding agent to compete with a WW domain substrate for binding to a WW domain, wherein the potential binding agent is modeled to fit spatially into a WW domain interaction site defined by a plurality of atomic coordinates.

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27. The crystalline complex of claim 23, wherein said binding agent is a C-terminal domain of RNA polymerase II.
28. The crystalline complex of claim 23, wherein said binding agent has a sequence Tyr-pSer-Pro-Thr-pSer-Pro-Ser (SEQ ID NO:3).